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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
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PERLEGEN SCIENCES, INC.
LEGAL DEPARTMENT
2021 STIERLIN COURT
MOUNTAIN VIEW, CA 94043

EXAMINER

FREDMAN, JEFFREY NORMAN

ART UNIT PAPER NUMBER

1637

DATE MAILED: 04/28/2005

Please find below and/or attached an Office communication concerning this application or proceeding.

Office Action Summary

Application No.

09/938,878

Applicant(s)

PATIL ET AL.

Examiner

Jeffrey Fredman

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-- The MAILING DATE of this communication appears on the cover sheet with the correspondence address --

Period for Reply

A SHORTENED STATUTORY PERIOD FOR REPLY IS SET TO EXPIRE 3 MONTH(S) FROM THE MAILING DATE OF THIS COMMUNICATION.

- Extensions of time may be available under the provisions of 37 CFR 1.136(a). In no event, however, may a reply be timely filed after SIX (6) MONTHS from the mailing date of this communication.
- If the period for reply specified above is less than thirty (30) days, a reply within the statutory minimum of thirty (30) days will be considered timely.
- If NO period for reply is specified above, the maximum statutory period will apply and will expire SIX (6) MONTHS from the mailing date of this communication.
- Failure to reply within the set or extended period for reply will, by statute, cause the application to become ABANDONED (35 U.S.C. § 133). Any reply received by the Office later than three months after the mailing date of this communication, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b).

Status

- 1) ☒ Responsive to communication(s) filed on 21 March 2005.
- 2a) ☐ This action is **FINAL**. 2b) ☒ This action is non-final.
- 3) ☐ Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under *Ex parte Quayle*, 1935 C.D. 11, 453 O.G. 213.

Disposition of Claims

- 4) ☒ Claim(s) 24-62 is/are pending in the application.
- 4a) Of the above claim(s) _____ is/are withdrawn from consideration.
- 5) ☐ Claim(s) _____ is/are allowed.
- 6) ☒ Claim(s) 24-62 is/are rejected.
- 7) ☐ Claim(s) _____ is/are objected to.
- 8) ☐ Claim(s) _____ are subject to restriction and/or election requirement.

Application Papers

- 9) ☐ The specification is objected to by the Examiner.
- 10) ☐ The drawing(s) filed on _____ is/are: a) ☐ accepted or b) ☐ objected to by the Examiner.
Applicant may not request that any objection to the drawing(s) be held in abeyance. See 37 CFR 1.85(a).
Replacement drawing sheet(s) including the correction is required if the drawing(s) is objected to. See 37 CFR 1.121(d).
- 11) ☐ The oath or declaration is objected to by the Examiner. Note the attached Office Action or form PTO-152.

Priority under 35 U.S.C. § 119

- 12) ☐ Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).
- a) ☐ All b) ☐ Some * c) ☐ None of:
1. ☐ Certified copies of the priority documents have been received.
2. ☐ Certified copies of the priority documents have been received in Application No. _____.
3. ☐ Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).
- * See the attached detailed Office action for a list of the certified copies not received.

Attachment(s)

- 1) ☐ Notice of References Cited (PTO-892)
- 2) ☐ Notice of Draftsperson's Patent Drawing Review (PTO-948)
- 3) ☒ Information Disclosure Statement(s) (PTO-1449 or PTO/SB/08)
Paper No(s)/Mail Date 9/12/04, 3/2/05
- 4) ☐ Interview Summary (PTO-413)
Paper No(s)/Mail Date. _____
- 5) ☐ Notice of Informal Patent Application (PTO-152)
- 6) ☐ Other: _____

DETAILED ACTION

Continued Examination Under 37 CFR 1.114

1. A request for continued examination under 37 CFR 1.114, including the fee set forth in 37 CFR 1.17(e), was filed in this application after final rejection. Since this application is eligible for continued examination under 37 CFR 1.114, and the fee set forth in 37 CFR 1.17(e) has been timely paid, the finality of the previous Office action has been withdrawn pursuant to 37 CFR 1.114. Applicant's submission filed on March 17, 2005 has been entered.

Claim Rejections - 35 USC § 103

2. The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:

(a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negated by the manner in which the invention was made.

3. This application currently names joint inventors. In considering patentability of the claims under 35 U.S.C. 103(a), the examiner presumes that the subject matter of the various claims was commonly owned at the time any inventions covered therein were made absent any evidence to the contrary. Applicant is advised of the obligation under 37 CFR 1.56 to point out the inventor and invention dates of each claim that was not commonly owned at the time a later invention was made in order for the examiner to consider the applicability of 35 U.S.C. 103(c) and potential 35 U.S.C. 102(e), (f) or (g) prior art under 35 U.S.C. 103(a).

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4. Claims 24, 29, 31, 38-40 are rejected under 35 U.S.C. 103(a) as being unpatentable over Zonana et al (U.S. Patent 6,355,782) in view of Dong et al (U.S. Patent 6,361,947).

Zonana teaches a method of analyzing a subset of nucleic acids (see abstract) comprising:

(a) providing a driver population of nucleic acid and a tester population of nucleic acids (See column 22, lines 64-65 for driver and column 22, line 66 to column 23, line 8 for tester),

(b) denaturing said population of tester and driver nucleic acids (see column 23, lines 9-16),

(c) annealing the driver and tester populations to produce a single stranded subset of nucleic acids and a double stranded subset of nucleic acids (see column 23, lines 15-18),

(d) immobilizing the driver population of nucleic acids by use of a biotin-streptavidin interaction to produce an unimmobilized single stranded tester subset of nucleic acids, an immobilized double stranded tester-driver subset of nucleic acids and an immobilized single stranded driver subset of nucleic acids (see column 23, lines 18-19),

(e) separating the unimmobilized single stranded tester subset of the nucleic acids from the single and double stranded driver subset of the nucleic acids (see column 23, lines 20-21),

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(f) dissociating the immobilized double stranded tester-driver subset of nucleic acids to produce a subset of complementary tester nucleic acids and a subset of immobilized complementary driver nucleic acids (see column 23, lines 22-23)

(g) separating the subset of complementary tester nucleic acids from the subset of immobilized complementary driver nucleic acids (see column 23, lines 22-23),

Zonana does not teach the steps of:

(h) hybridizing the unimmobilized single stranded tester nucleic acids to probes on a nucleic acid probe array (see page 1890, column 2, subheading "colony hybridization" and figure 3) and

(i) determining which of the probes on the array hybridizes to the single stranded tester subset of the population thereby analyzing the single stranded subset of the population of nucleic acid fragments (see page 1890, column 2, subheading "colony hybridization and figure 3).

Dong teaches the steps of

(h) hybridizing the unimmobilized single stranded tester nucleic acids to probes on a nucleic acid probe array (see column 5, lines 57-60 and column 31, claim 1)) where Dong further teaches that "In a preferred embodiment the isolated sequences are then exposed to an array which may or may not have been specifically designed and manufactured to interrogate the isolated sequences. Design of both the complexity management steps and the arrays may be aided by the computer modeling techniques which are also described in the present invention (see column 5, lines 58-61)" and

(i) determining which of the probes on the array hybridizes to the single stranded tester subset of the population thereby analyzing the single stranded subset of the population of nucleic acid fragments (see column 5, lines 57-60 and column 31, claim 1).

With regard to claim 29, Zonana teaches the use of PCR products as driver (see column 22, lines 64-65).

With regard to claims 38 and 39, Zonana teaches the driver has a biotin tag and binds to streptavidin magnetic beads (see column 23, lines 18-19).

With regard to claim 40, Zonana teaches separating the subset of complementary tester nucleic acids from the subset of immobilized complementary driver nucleic acids using the biotin streptavidin interaction (see column 23, lines 22-23),

With regard to claim 31, Dong teaches restriction digestion of the sample which will result in more than ten noncontiguous regions of driver (see columns 6-8).

It would have been prima facie obvious to one of ordinary skill in the art at the time the invention was made to combine the method of Zonana with the detection method of Dong since Zonana wants to selected cDNA and since Dong states "In a preferred embodiment the isolated sequences are then exposed to an array which may or may not have been specifically designed and manufactured to interrogate the isolated sequences. (see column 5, lines 57-60)." An ordinary practitioner would have recognized that both Zonana and Dong were operating to reduce the complexity of their DNA sample and were selecting for subsets of the total sample. In this context, an ordinary practitioner would have been motivated by Dong to use an array in the

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place of the more cumbersome cloning methods used by Zonana for further analysis since Dong expressly teaches that array detection is a preferred method of analysis of the isolated subsets.

5. Claims 25-28, 30-37 and 41-62 are rejected under 35 U.S.C. 103(a) as being unpatentable over Zonana et al (U.S. Patent 6,355,782) in view of Dong et al (U.S. Patent 6,361,947) as applied to claims 24, 29 and 38-40 and further in view of Wigler et al (U.S. Patent 5,501,964).

Zonana et al (U.S. Patent 6,355,782) in view of Dong et al (U.S. Patent 6,361,947) teach the limitations of claims 24, 29 and 38-40 as discussed above. Zonana et al (U.S. Patent 6,355,782) in view of Dong et al (U.S. Patent 6,361,947) do not teach screening fragments from human individuals or the use of two different human individuals or comparison of different species or the DNA or mRNA sources used.

Wigler teaches comparison of DNA from two sources in order to determine the relationship between the sources (See column 3, lines 11-14) including comparisons between different individuals (see column 8, lines 40-48) as well as comparisons between different species (see column 21, example 7), which address the limitations of claims 27-28, 42, 49-50, 56-57.

With regard to claims 25-26, 46-48, 54-55, Wigler teaches that the sources can be cDNA, genomic DNA, restriction fragments of DNA or libraries (see column 2, lines 42-50).

With regard to claims 31, 52, 53, the cDNA drivers would necessarily be derived from noncontiguous regions of a genome of a species. Wigler also teaches comparison

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of PCR amplified DNA (see column 4, lines 28-37). Wigler expressly recognizes that any animal can be the source of the DNA, including mammals and non-mammals, as well as higher eukaryotes and humans.(see column 3, lines 62-67).

With regard to claims 41, 45, 51, 58, Zonana teaches the use of PCR products as driver prior to step (a) (see column 22, lines 64-65) and Wigler teaches the use of genomic DNA (see column 2, lines 42-50).

With regard to claim 43, Zonana teaches the use of PCR and no specific length difference exists for "long range PCR" in the specification and so this term does not distinguish from ordinary PCR.

With regard to claim 44, 51, Zonana teaches PCR to amplify tester nucleic acid (see column 23, lines 1-15).

With regard to claims 60-61, Zonana teaches the driver has a biotin tag and binds to streptavidin magnetic beads (see column 23, lines 18-19).

With regard to claim 62, Zonana teaches immobilizing the driver population of nucleic acids by use of a biotin-streptavidin interaction to produce an unimmobilized single stranded tester subset of nucleic acids, an immobilized double stranded tester-driver subset of nucleic acids and an immobilized single stranded driver subset of nucleic acids (see column 23, lines 18-19),

It would have been prima facie obvious to one of ordinary skill in the art at the time the invention was made to modify the method of Zonana et al (U.S. Patent 6,355,782) in view of Dong et al (U.S. Patent 6,361,947) to utilize the different comparisons and DNA sources for comparison taught by Wigler since Wigler states

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"Comparative genomic DNA analysis holds promise for the discovery of sequences which may provide for information concerning polymorphisms, infectious DNA based agents, lesions associated with disease, such as cancer, inherited dominant and recessive traits, and the like. By being able to detect particular DNA sequences which have a function or affect a function of cells, one can monitor pedigrees, so that in breeding animals one can follow the inheritance of particular sequences associated with desirable traits. In humans, there is substantial interest in forensic medicine, diagnostics and genotyping, and determining relationships between various individuals. There is, therefore, substantial interest in providing techniques which allow for the detection of common sequences between sources and sequences which differ between sources. (Column 1, lines 23-37)."

An ordinary practitioner would have been motivated to apply the tester driver method of Zonana et al (U.S. Patent 6,355,782) in view of Dong et al (U.S. Patent 6,361,947) on comparisons between individuals and between species in order to identify desirable traits, as expressly suggested by Wigler, as well as identifying relationships between individuals and species as suggested by Wigler. An ordinary practitioner would have been motivated to focus on a comparison of unique sequences as taught by Straus in the broad variety of contexts suggested by Wigler.

Further, with regard to the order of the steps of immobilization, annealing and denaturation, as in claim 59, for example, as MPEP 2144.04 notes "selection of any order of performing process steps is prima facie obvious in the absence of new or unexpected results". In this case, this is particularly true since the order of the steps would not be expected to impact the results of the method. Whether immobilization was performed prior to the annealing or denaturation steps would not be expected to effect the reaction since the interaction is between the nucleic acids, which are equally available whether immobilized or not.

Response to Arguments


6. There are no new arguments, and the rejection has been altered to address the new claim limitations.

Conclusion

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Jeffrey Fredman whose telephone number is (571)272-0742. The examiner can normally be reached on 6:30-3:00.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Gary Benzion can be reached on (571)272-0782. The fax phone number for the organization where this application or proceeding is assigned is 703-872-9306.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see <http://pair-direct.uspto.gov>. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).


Jeffrey Fredman
Primary Examiner
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